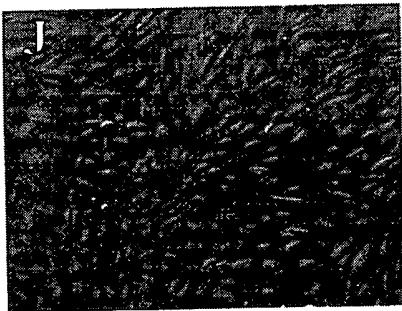
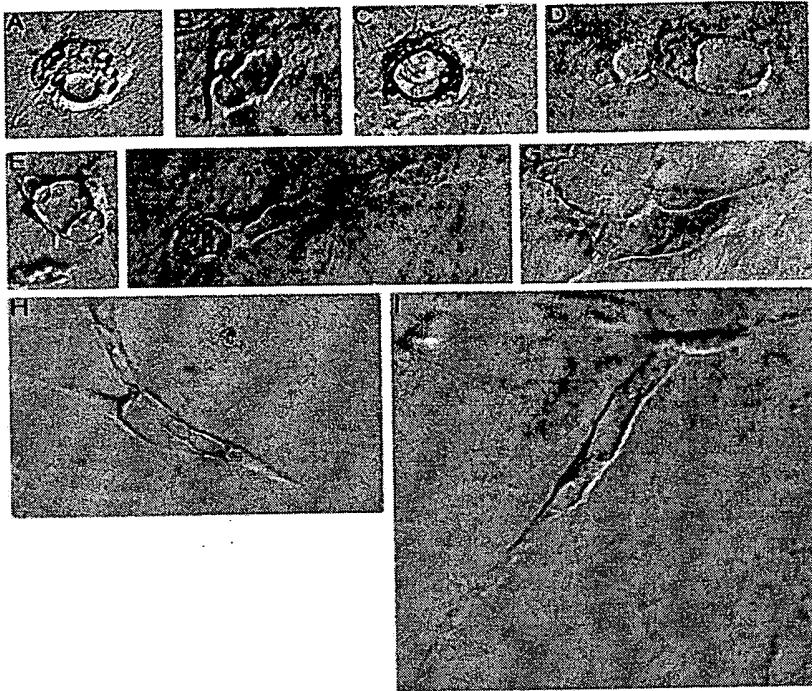


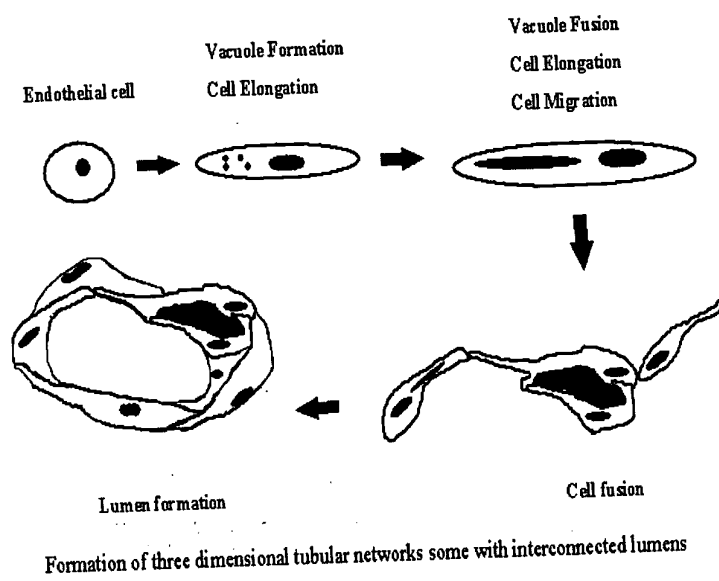
**FIGURE 1**



HUVEC on collagen film, 48 hr.

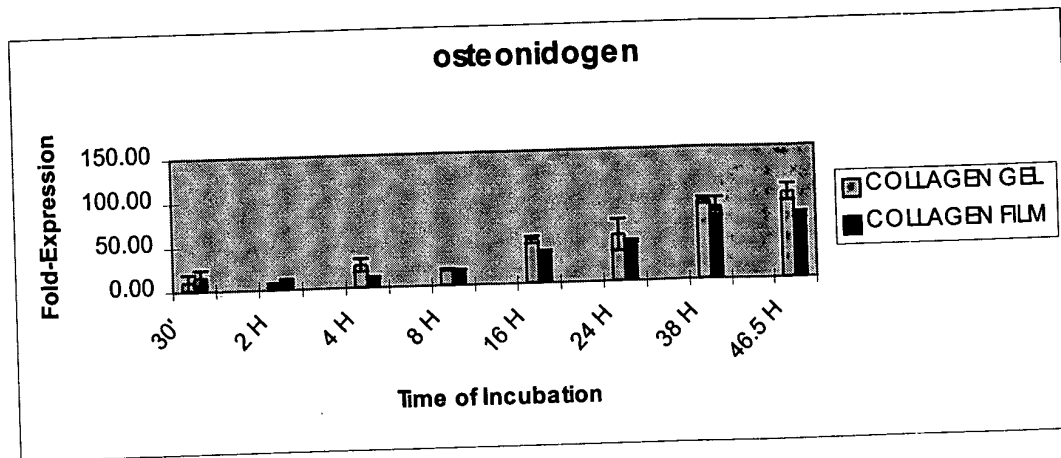
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**FIGURE 2**

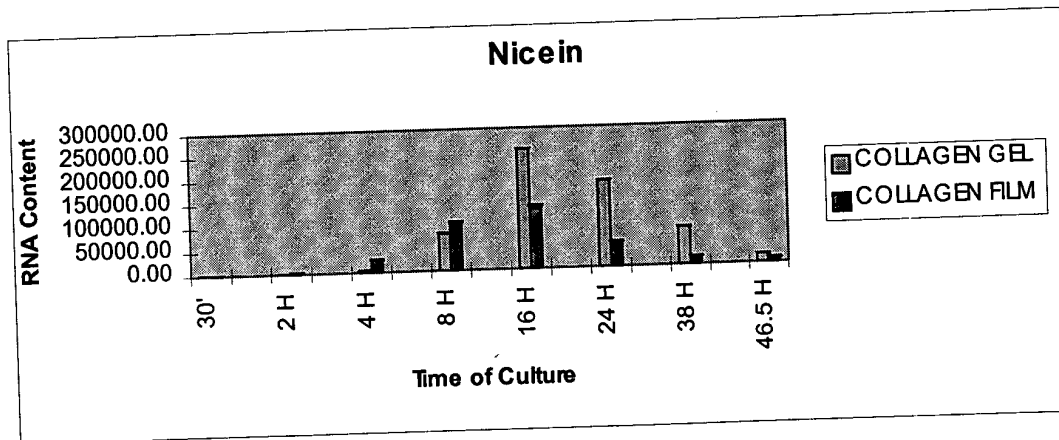


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**FIGURE 3: PA1 (osteonidogen)**

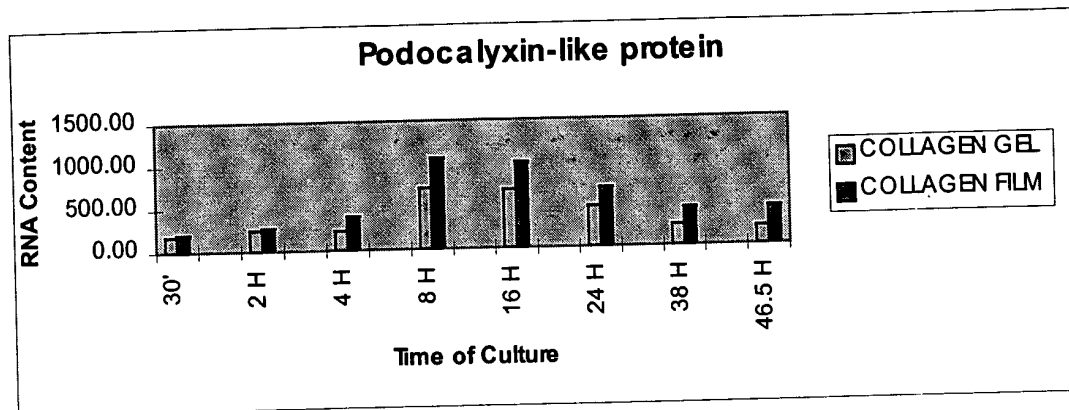


**FIGURE 4: PA2 (laminin gamma-2 chain (nicein))**

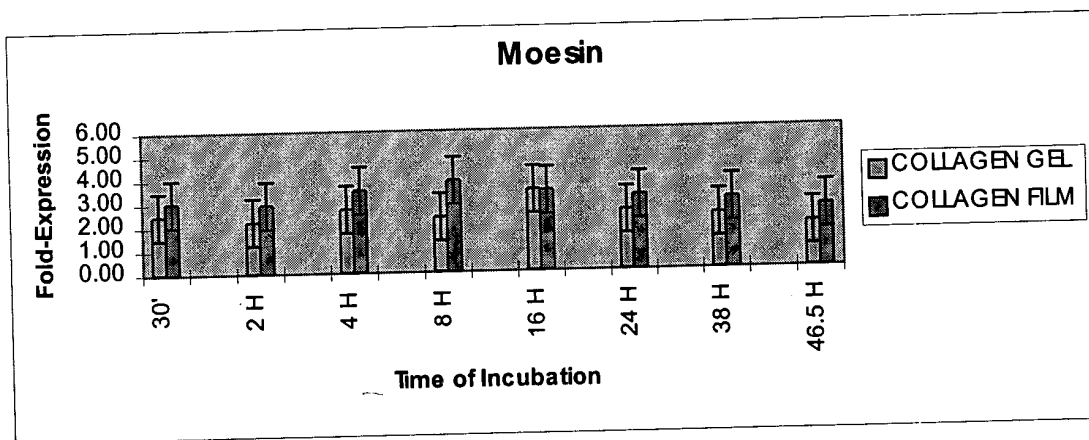


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**FIGURE 5: PA3 (podocalyxin like protein)**

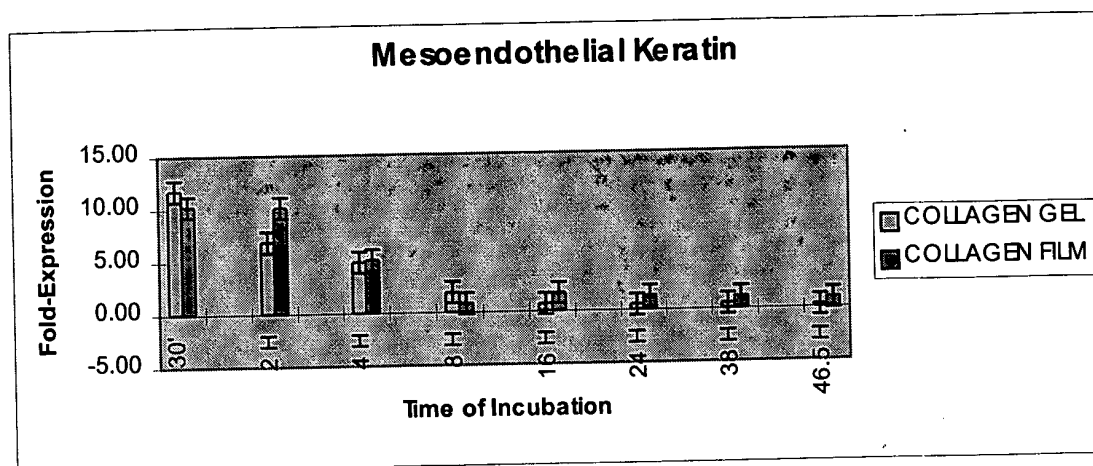


**FIGURE 6: PA4 (moesin)**

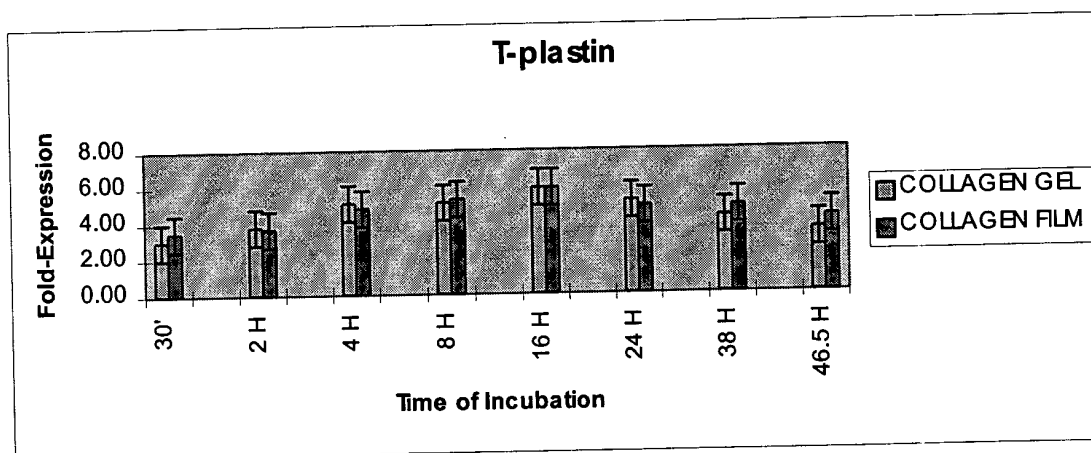


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**FIGURE 7: PA5 (mesoendothelial keratin K7)**

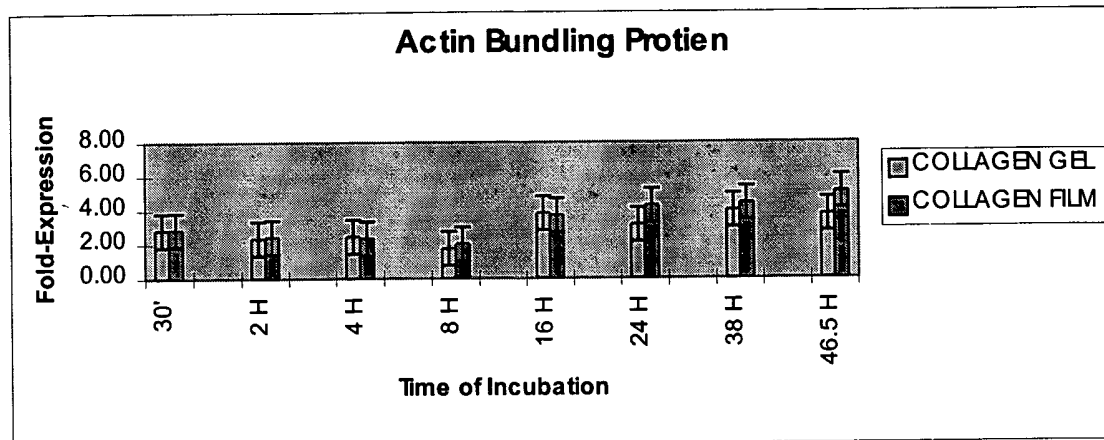


**FIGURE 8: PA7 (T-plastin)**

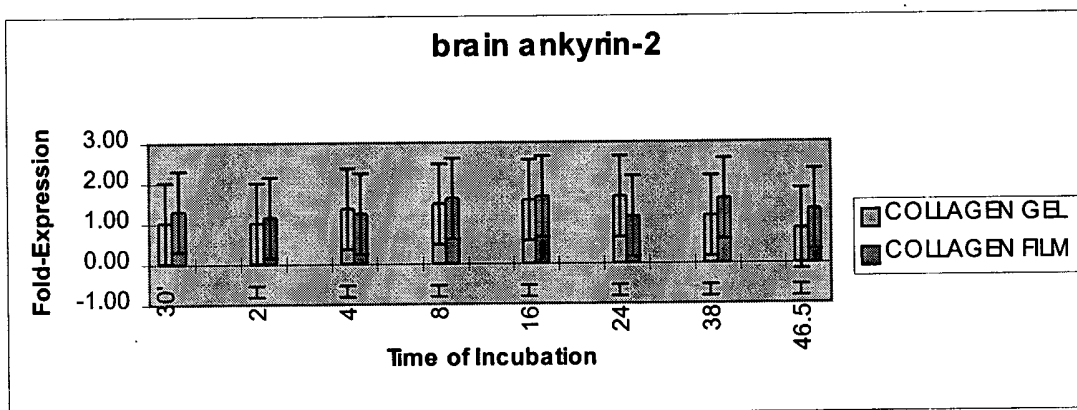


DOT-01-05-00/60

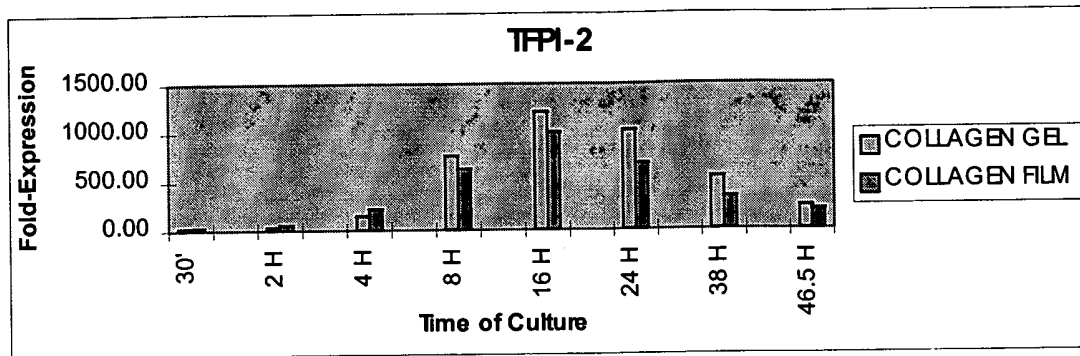
**FIGURE 9: PA8 (actin bundling protein)**



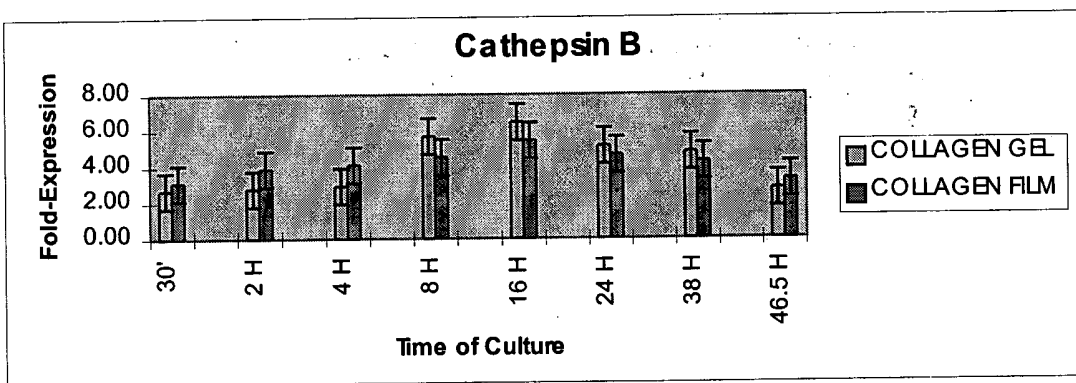
**FIGURE 10: PA10 (C3VS homolog)**



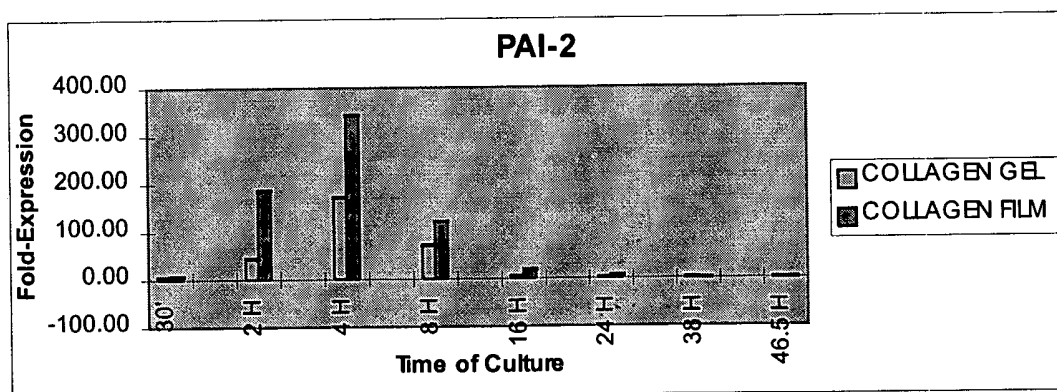
**FIGURE 11: PA13 (tissue factor pathway inhibitor)**



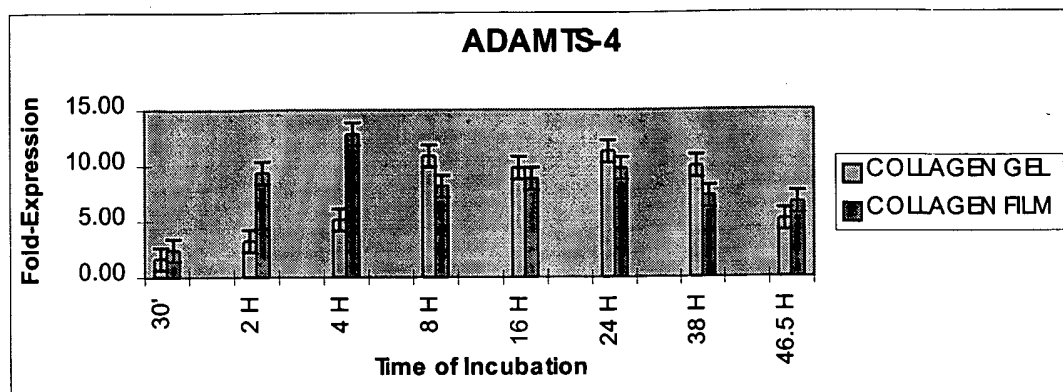
**FIGURE 12: PA11 (cathepsin B)**



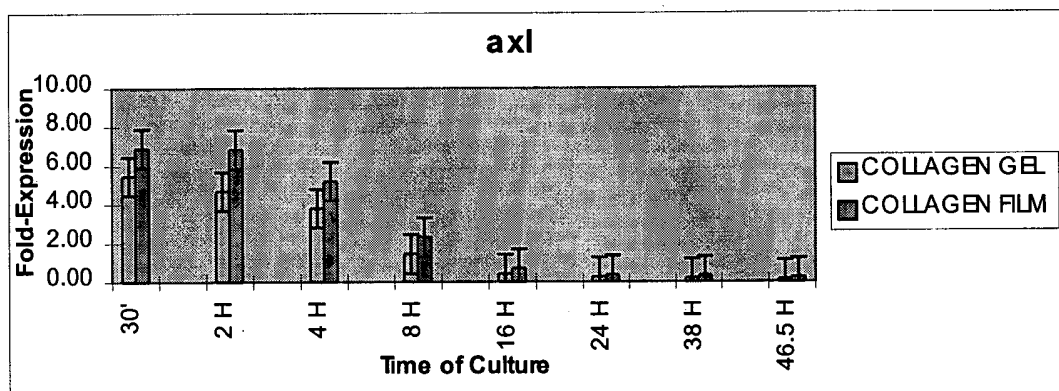
**FIGURE 13: PA14 (plasminogen activator inhibitor)**



**FIGURE 14: PA12 (ADAMTS-4)**

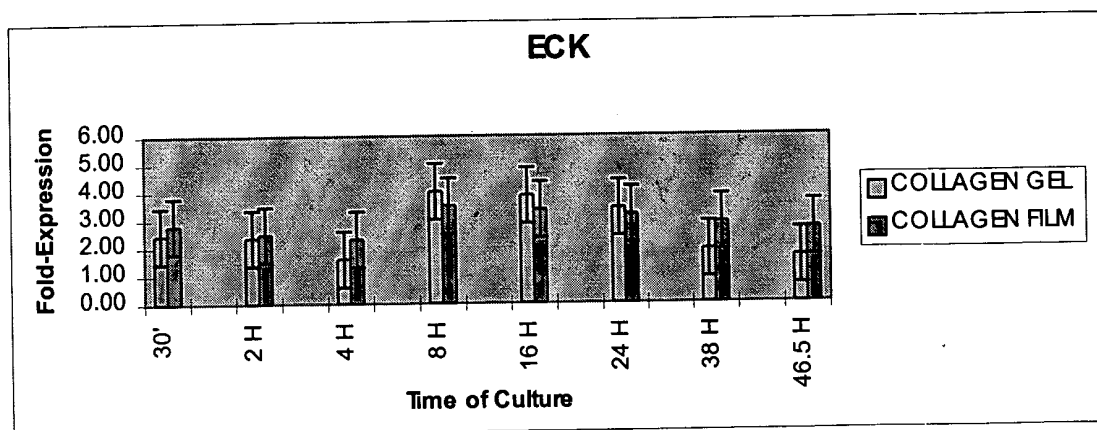


**FIGURE 15: PA15 (axl)**

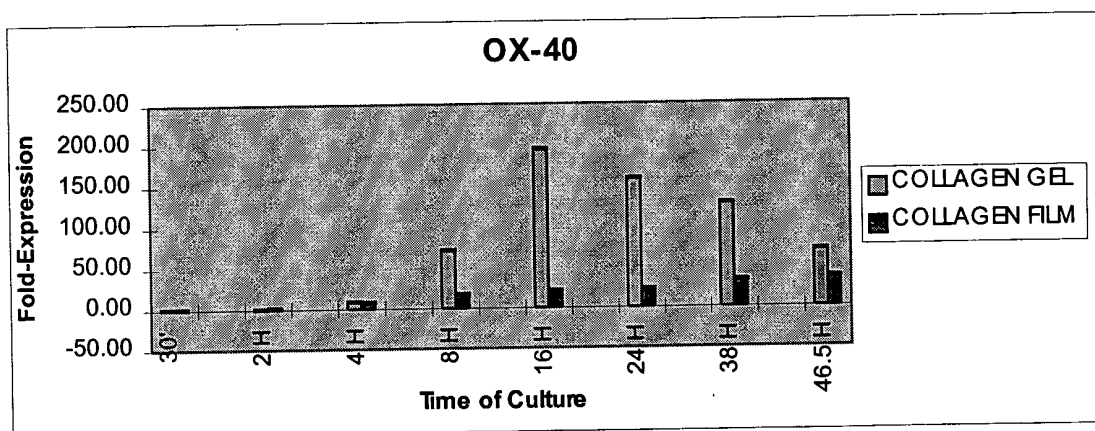




**FIGURE 16: PA16 (ECK)**

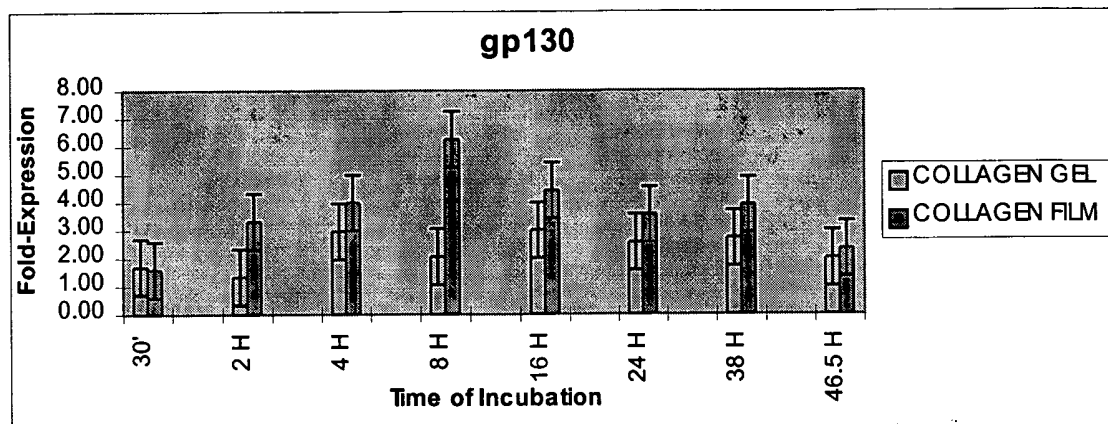


**FIGURE 17: PA17 (OX-40)**

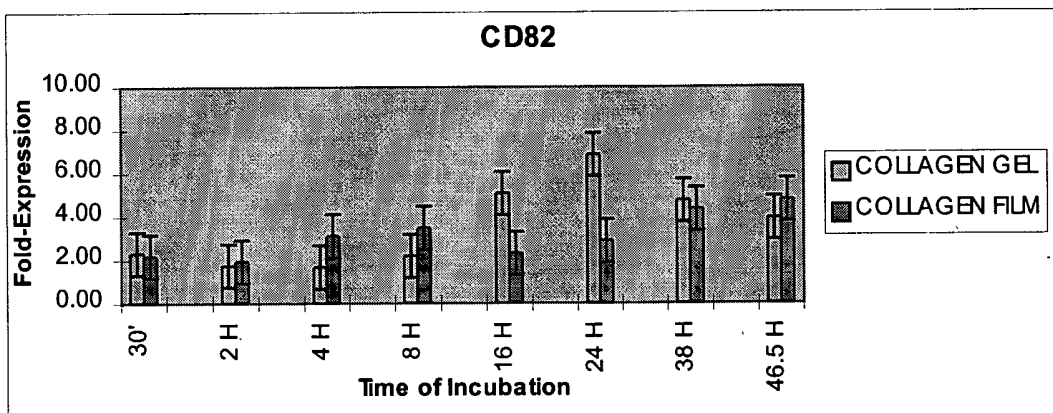


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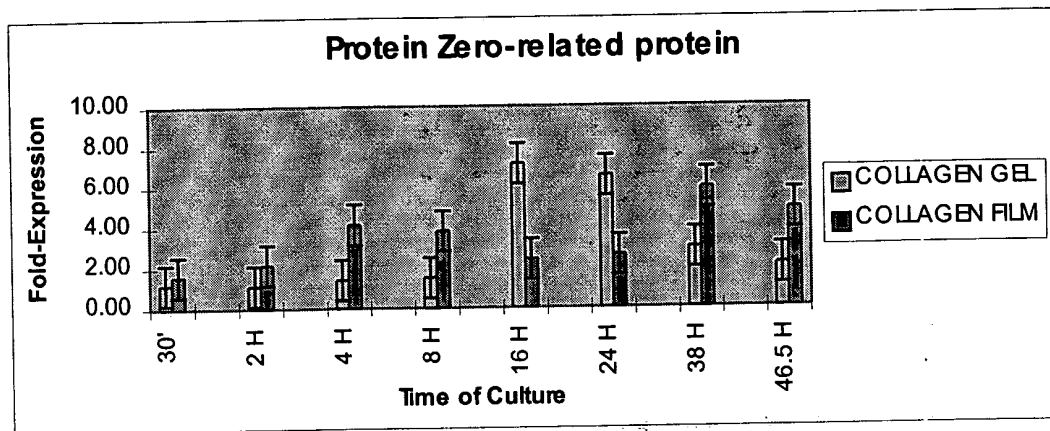
**FIGURE 18: PA18 (gp130)**



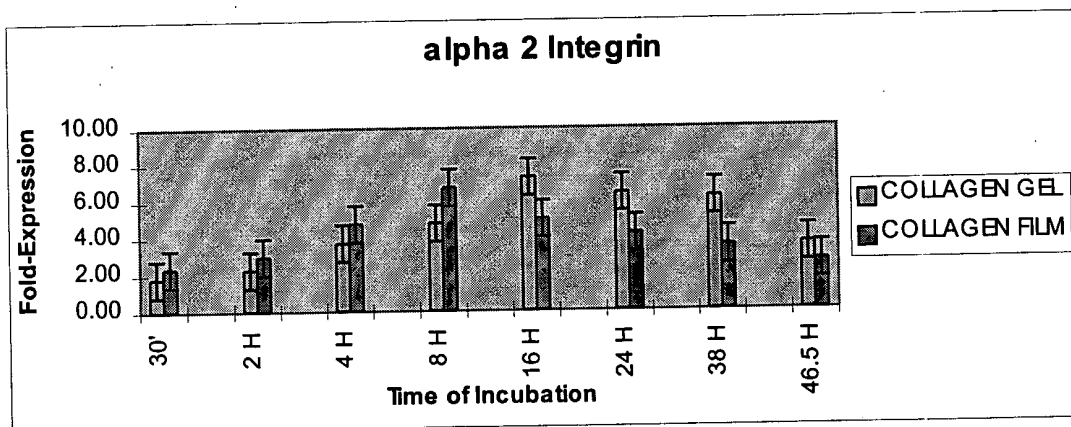
**FIGURE 19: PA19 (CD82)**



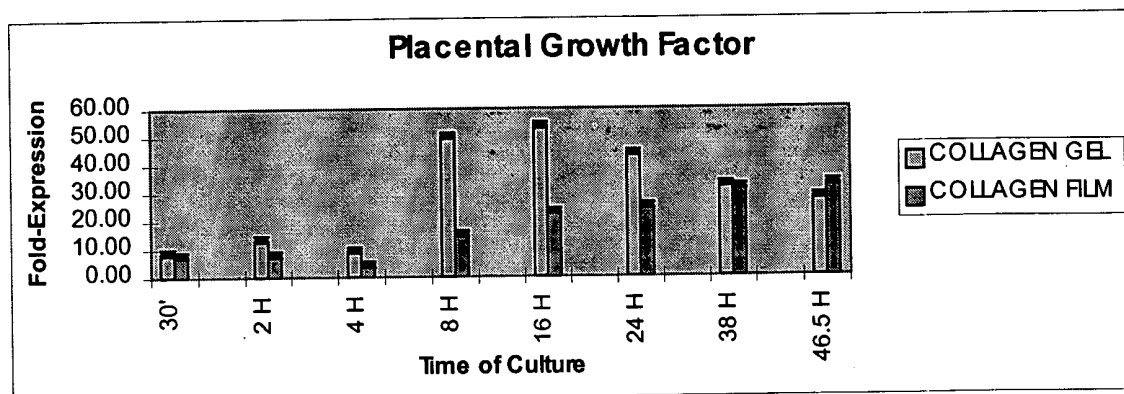
**FIGURE 20: PA20 (PRZ)**



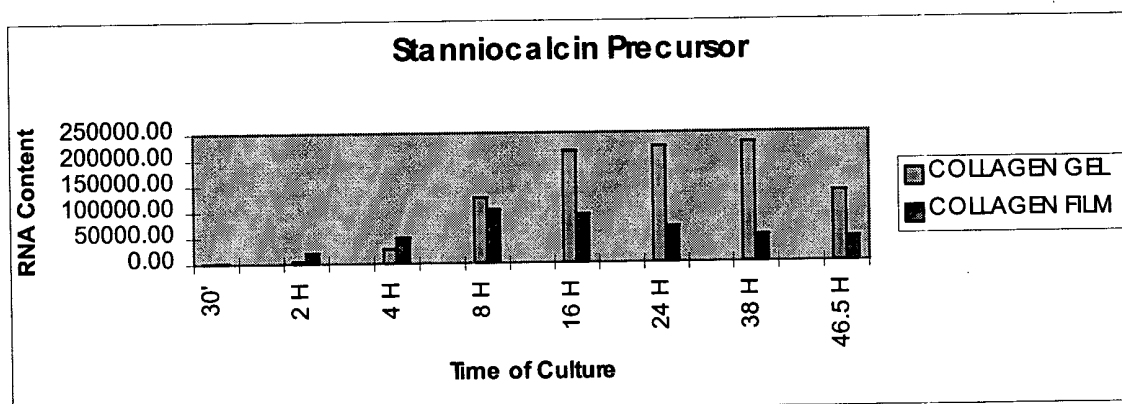
**FIGURE 21: PA21 (alpha 2 integrin)**



**FIGURE 22: PA22 (PIGF)**

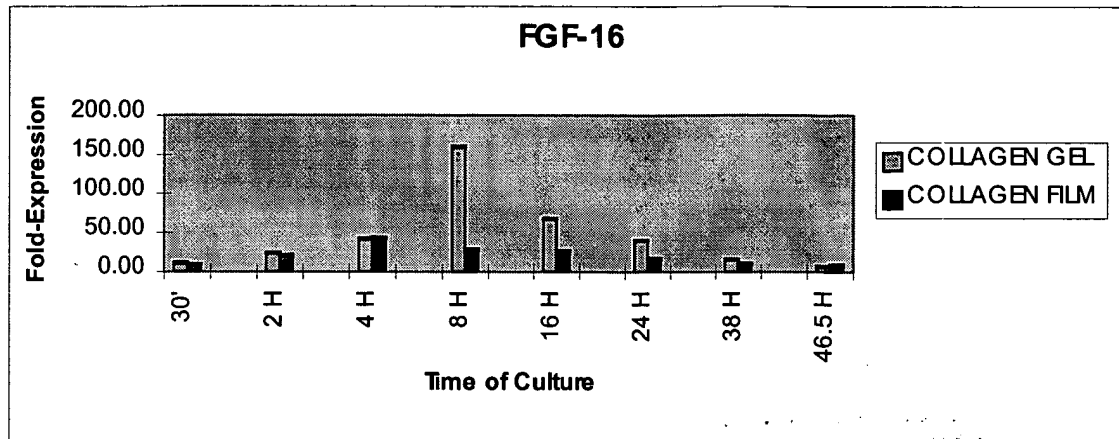


**FIGURE 23: PA23 (stanniocalcin precursor)**



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**FIGURE 24: PA24 (FGF-16)**



**FIGURE 25: PA25 (White Protein Homolog)**

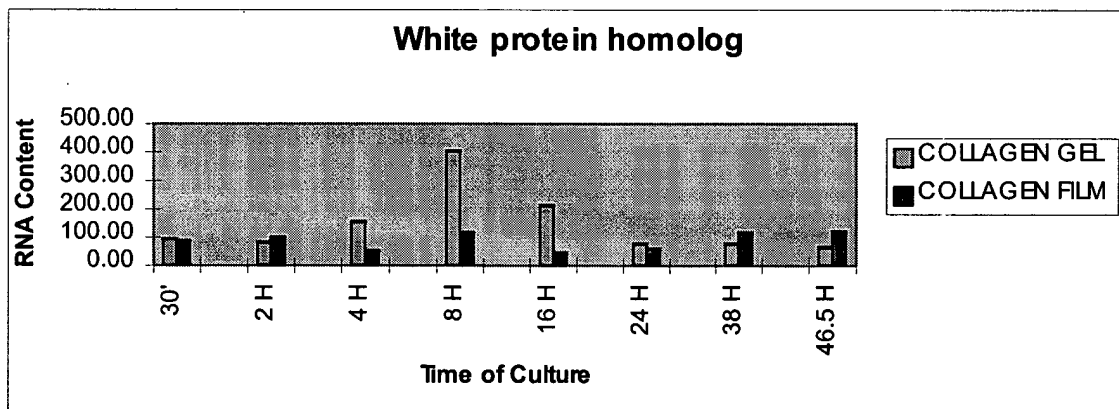
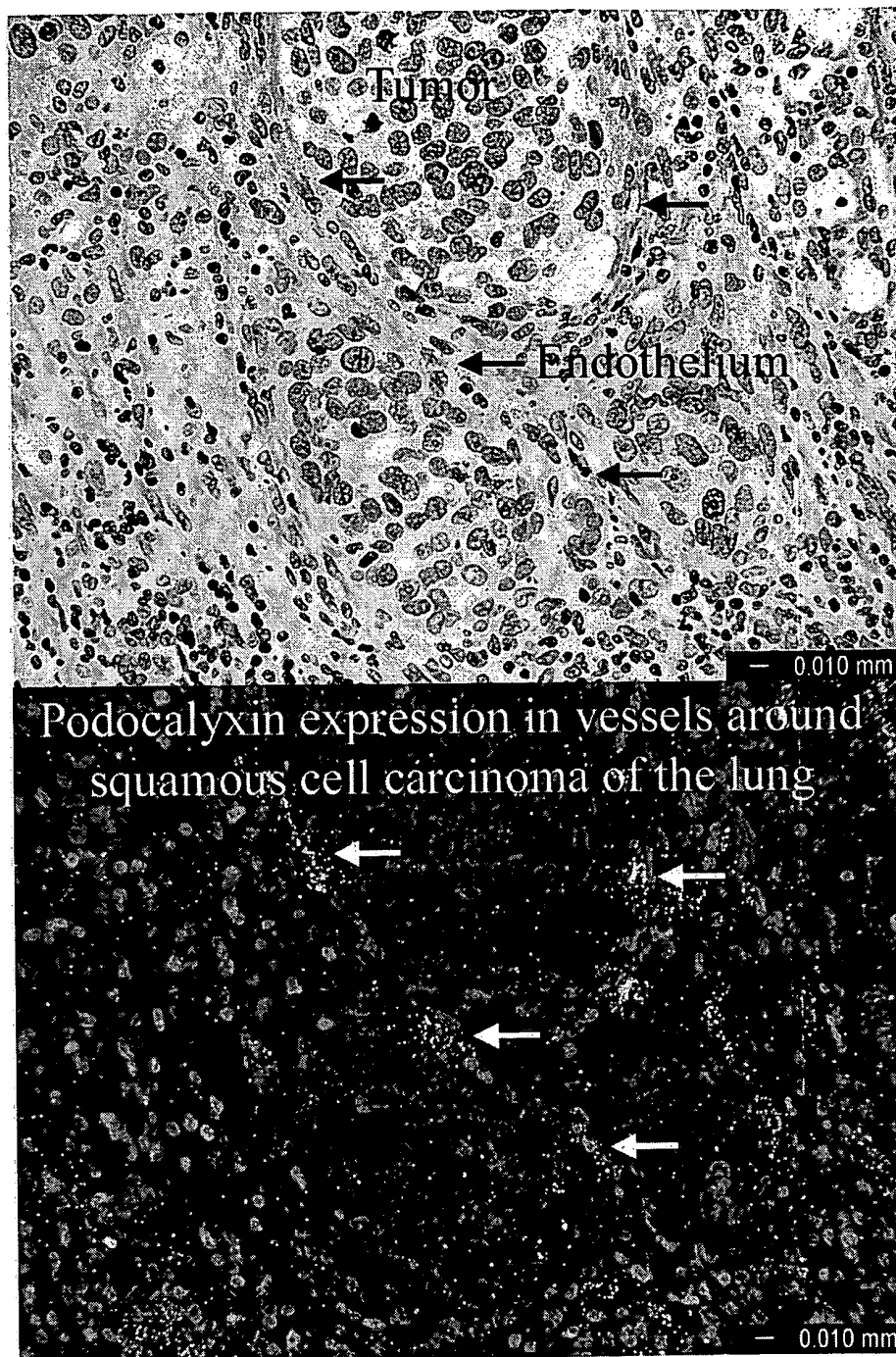


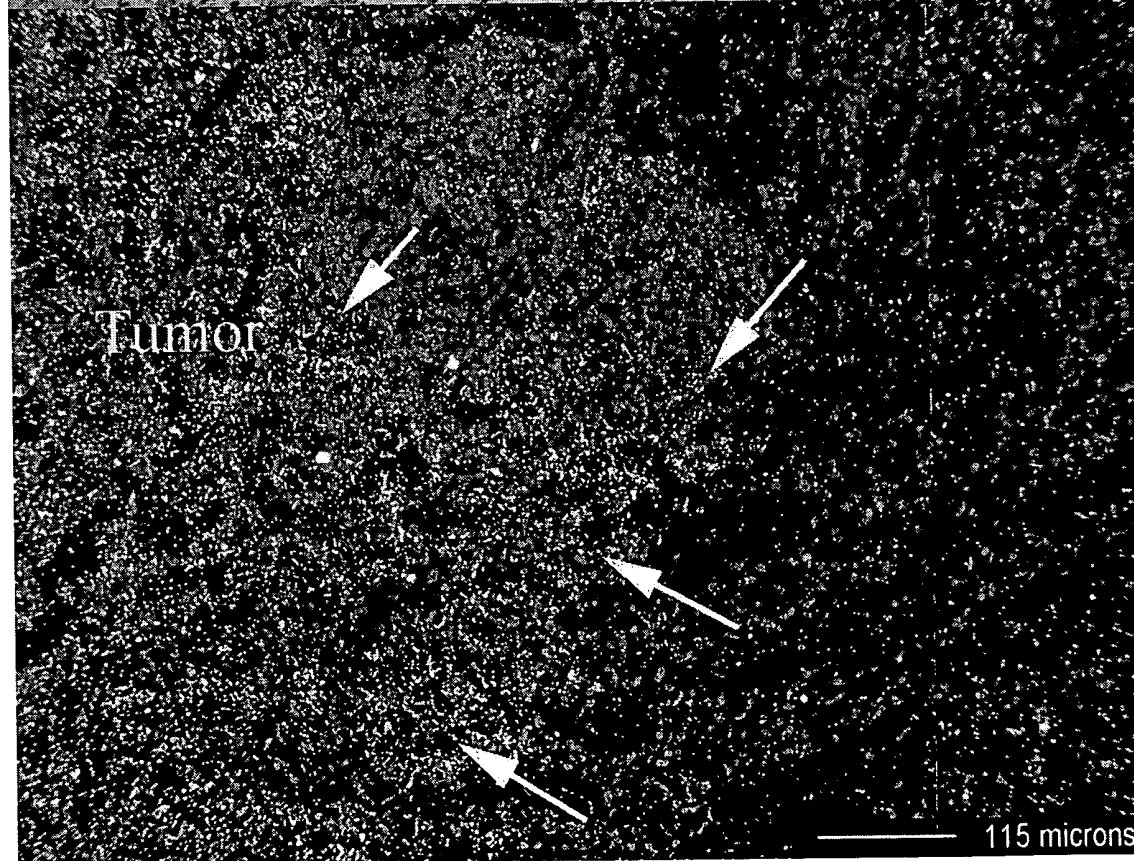
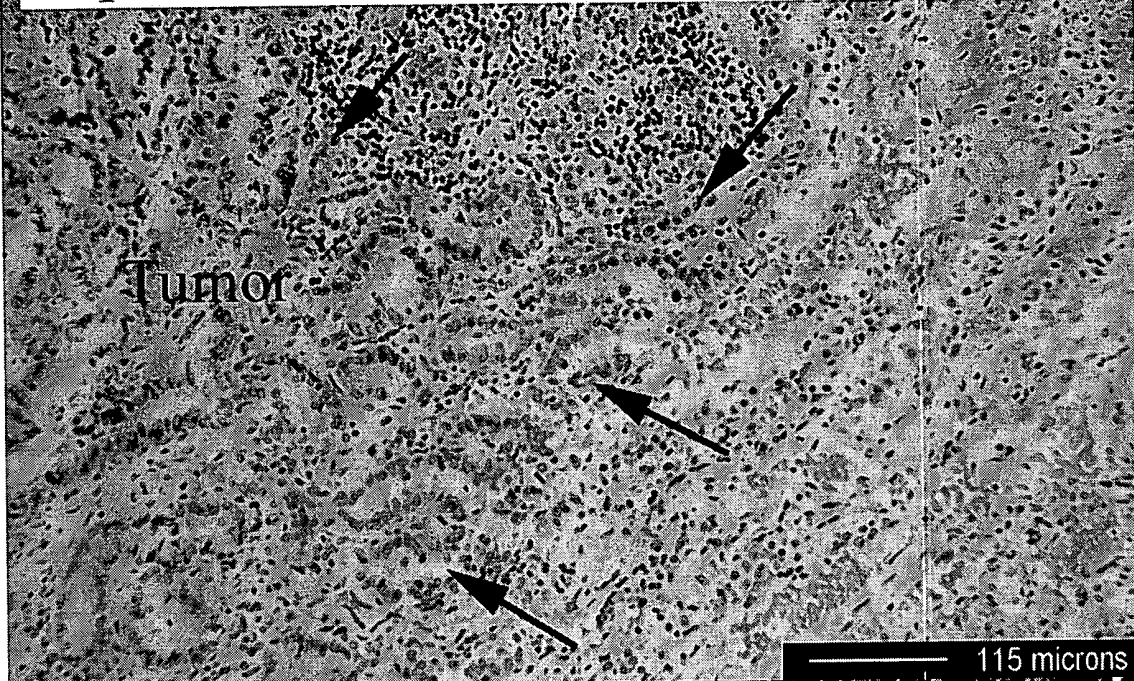
FIGURE 26



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FIGURE 27.

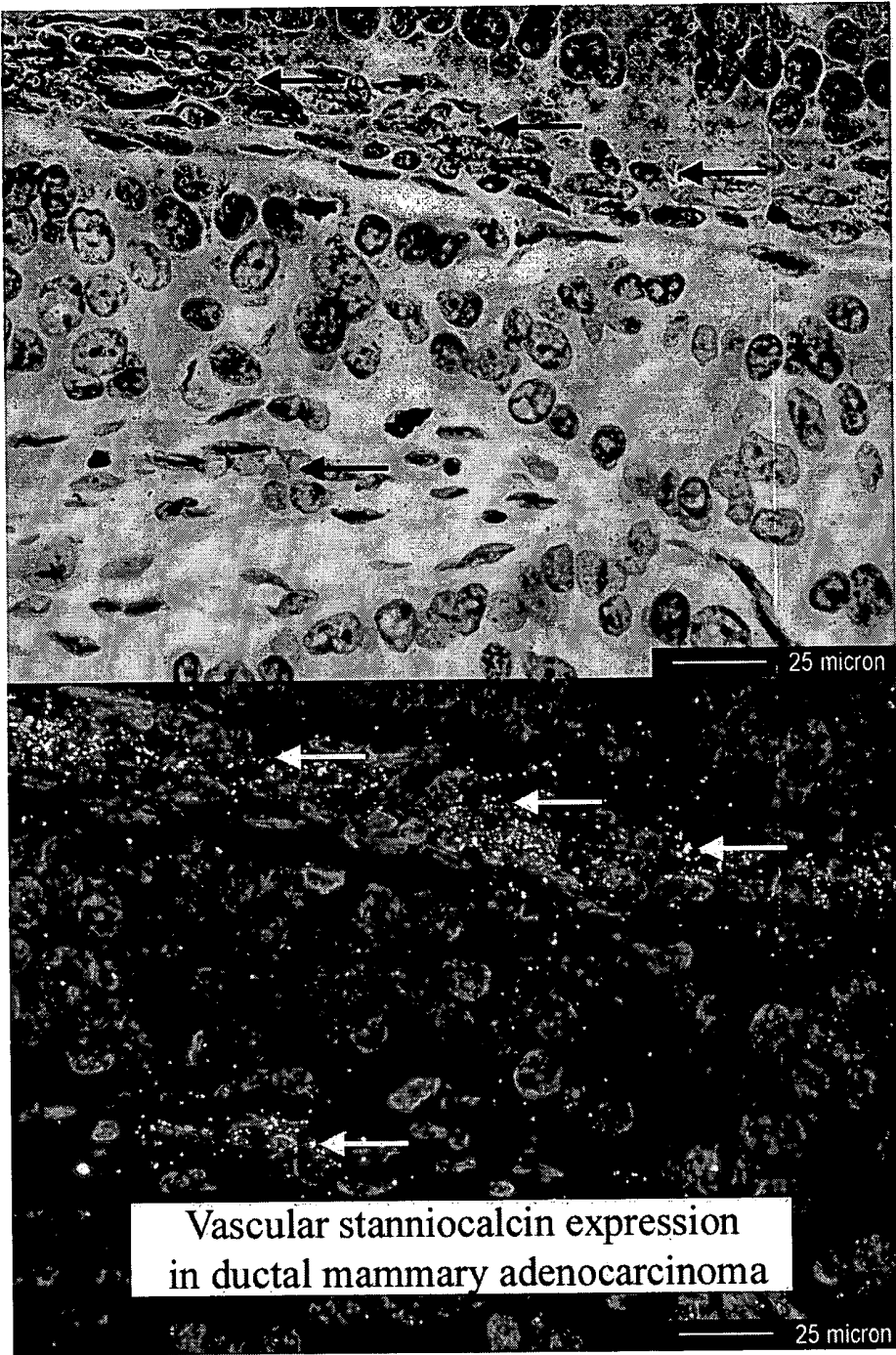
# DNA 76510 (Protein Zero Homolog) Expression in Pulmonary Adenocarcinoma



0070350-10100



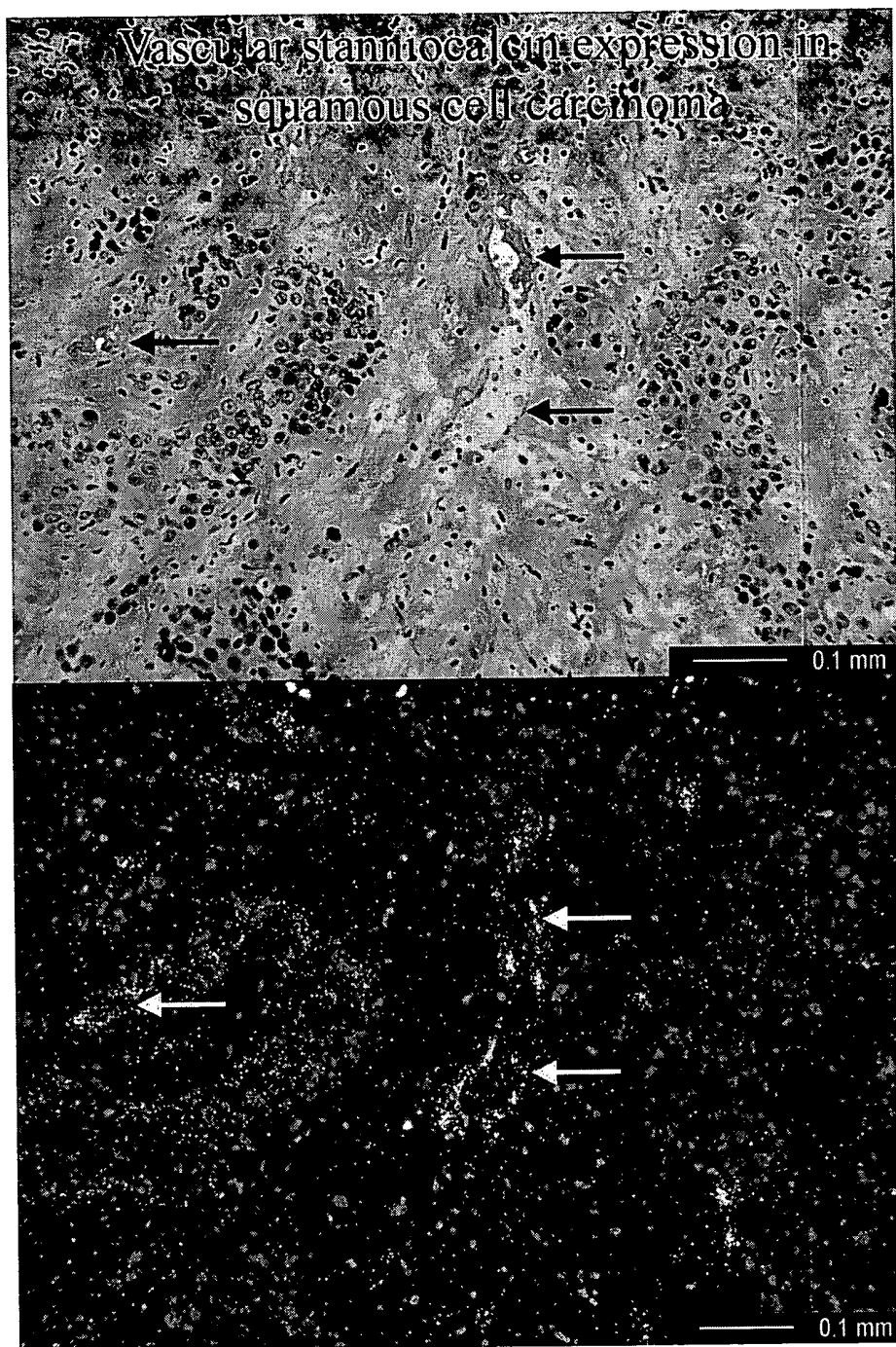
FIGURE 28



00703350-10100



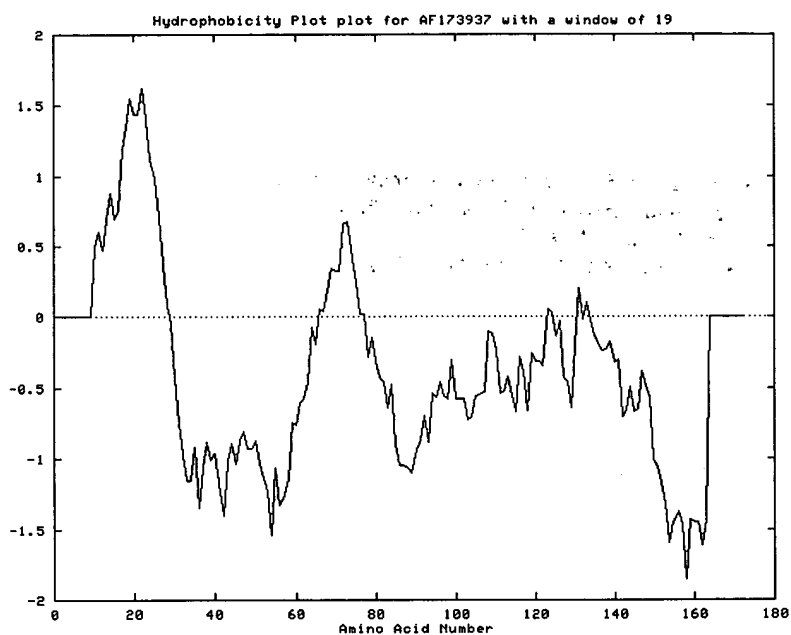
FIGURE 29



**FIGURE 30**

1 HVGPAPRRRLRPLAALALVLALAPGLPTARAGQTPRPAEROPPVRLLFTEEELARYGGELEDQPIYLAVKGVVFDVTSCKE  
01 FYGRGAPYNALTGRDSTRGVAKHSLDPADLTHDTTGLTAKELEALDEVFTKVYKAKYPIVGYTARRILNEDGSPNLDPKP  
161 EDQPHFDIKDEF (SEQ. ID. NO.: 72)

**FIGURE 31**



05703350-103100

**FIGURE 32**

Sequence type explicitly set to Protein  
Sequence format is Pearson  
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Sequence 2: 11725372\_EXT 161 aa  
Sequence 3: AF173937 172 aa  
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Aligning...  
Sequences (1:2) Aligned. Score: 83  
Sequences (1:3) Aligned. Score: 83  
Sequences (2:3) Aligned. Score: 90  
Start of Multiple Alignment  
There are 2 groups  
Aligning...  
Group 1: Sequences: 2 Score:1860  
Group 2: Sequences: 3 Score:1948  
Alignment Score 2458  
CLUSTAL-Alignment file created [/data4/genetools/lrastelli5501clustalw]

**Multiple Alignment:**

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AF173937      MVGPAPRRRLRPLAALALVLAAPGLEFARAGCTPRPAERGPPVRLFTEELARYGEEEE

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